



SEQUENCE LISTING

<110> Wadhwa, Renu
Sugihara, Takashi
Ohide, Akiko

<120> TUMOR SUPPRESSOR GENE

<130> 06501-091001

<140> US 10/045,815
<141> 2001-10-26

<150> PCT/JP00/02731
<151> 2000-04-26

<150> JP 11/118806
<151> 1999-04-26

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<170> FastSEQ for Windows Version 4.0

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Pro Asp Leu Leu Phe Ala Glu Gly Thr Ala Ala Tyr Ala Arg Gly Asp	
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Trp Pro Gly Val Val Leu Ser Met Glu Arg Ala Leu Arg Ser Arg Ala	
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Ala Leu Arg Ala Leu Arg Leu Arg Cys Arg Thr Gln Cys Ala Ala Asp	
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<212> PRT

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Lys	Ala	Val	Ala	Ala	Ala	His	Thr	Phe	Phe	Val	Gly	Asn	Pro	Glu	His
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Glu	Cys	Arg	Ala	Leu	Cys	Glu	Gly	Pro	Tyr	Asp	Tyr	Asp	Gly	Tyr	Asn
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105

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gtt gct gca gca cac acc ttc ttc gtg ggc aat cct gag cac atg gaa Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro Glu His Met Glu 165 170 175			585
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Cys Gln Glu Leu Gln Arg Leu Thr Asn Val Ala Ala Thr Ser Gly Asp	
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ggc tac cgg ggt cag acc tcc cca cat act ccc aat gaa aag ttc tat	1593
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Tyr Ser His Leu Val Cys Arg Thr Ala Ile Glu Glu Val Gln Ala Glu	
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Arg Lys Asp Asp Ser His Pro Val His Val Asp Asn Cys Ile Leu Asn	
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Ala Glu Thr Leu Val Cys Val Lys Glu Pro Pro Ala Tyr Thr Phe Arg	
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Asp Tyr Ser Ala Ile Leu Tyr Leu Asn Gly Asp Phe Asp Gly Gly Asn	
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Pro Gln Cys Gly Arg Ala Val Gly Phe Ser Ser Gly Thr Glu Asn Pro	
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His Gly Val Lys Ala Val Thr Arg Gly Gln Arg Cys Ala Ile Ala Leu	
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Trp Phe Thr Leu Asp Pro Arg His Ser Glu Arg Asp Arg Val Gln Ala	
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<212> PRT

<213> Homo sapiens

<400> 4

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Ala	Asp	Phe	Pro	Trp	Glu	Leu	Asp	Pro	Asp	Trp	Ser	Pro	Ser	Pro	Ala		
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Gln	Ala	Ser	Gly	Ala	Gly	Ala	Leu	Arg	Asp	Leu	Ser	Phe	Phe	Gly	Gly		
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			260					265					270			
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305					310					315					320	
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Ala	Glu	Arg	Lys	Asp	Asp	Ser	His	Pro	Val	His	Val	Asp	Asn	Cys	Ile	
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 Gly Arg Arg Arg Arg Arg Leu Thr Gly Cys Gly Glu Ser Glu Pro Gly
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tgg gac gtg gca gcc cct gac ctg ctt tac gca gag ggg acc gcg gcc 146
 Trp Asp Val Ala Ala Pro Asp Leu Leu Tyr Ala Glu Gly Thr Ala Ala
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 Tyr Ser Arg Arg Asp Trp Pro Gly Val Val Leu Asn Met Glu Arg Ala
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 Asp Pro Ser Leu Ser Gln Asp Pro Gly Ala Ala Ala Leu His Asp Leu
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Gly Asn Pro Glu His Met Glu Met Arg Gln Asn Leu Asp Tyr Tyr Gln	
175 180 185	
acc atg tct ggg gtg aag gag gca gac ttc agg gat ctc gag gcc aag	626
Thr Met Ser Gly Val Lys Glu Ala Asp Phe Arg Asp Leu Glu Ala Lys	
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Pro His Met His Glu Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu	
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Ile Thr Asp His Tyr Val Gln Val Leu Asn Cys Lys Gln Asn Cys Val	
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Thr Glu Leu Ala Ser His Pro Ser Arg Glu Lys Pro Phe Glu Asp Phe	
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Leu Pro Ser His Tyr Asn Tyr Leu Gln Phe Ala Tyr Tyr Asn Ile Gly	
305 310 315	
aac tat aca caa gct att gaa tgt gcc aag acc tac ctc ctc ttc ttt	1010
Asn Tyr Thr Gln Ala Ile Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe	
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Glu Tyr Arg Arg Pro Asn Leu Leu Glu Lys Glu Leu Leu Phe Phe Ala	
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Glu Thr Ala Val Arg Ile Ser Gln Glu Ile Gly Asn Leu Met Lys Glu	
415 420 425	
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Ile Glu Glu Ser Gln Ala Glu Arg Lys Asp Ser Ser His Pro Val His	
575 580 585	
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Val Asp Asn Cys Ile Leu Asn Ala Glu Ala Phe Met Cys Ile Lys Glu	
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Pro Pro Ala Tyr Thr Phe Arg Glu Tyr Ser Ala Ile Leu Tyr Leu Asn	

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tct tct ggc act gag aac cca cat gga gtg aag gct gtc acc agg ggg Ser Ser Gly Thr Glu Asn Pro His Gly Val Lys Ala Val Thr Arg Gly 655 660 665			2018
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ggc tgc cca gag cct gga gaa gag ttt ctg cat ggt gct act gtt ctt Gly Ser Pro Glu Pro Gly Glu Glu Phe Leu His Gly Ala Thr Val Leu 720 725 730			2210
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Asp	Glu	Glu	Cys	Arg	Ala	Leu	Cys	Glu	Gly	Pro	Tyr	Asp	Tyr	Asp	Gly
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Gln	Ala	Ile	Glu	Cys	Ala	Lys	Thr	Tyr	Leu	Leu	Phe	Phe	Pro	Asn	Asp
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 Thr Glu Leu Pro Trp Ala Pro Asp Leu Asp Leu Gly Pro Asp Pro Ser
 85 90 95
 Leu Ser Gln Asp Pro Gly Ala Ala Ala Leu His Asp Leu Arg Phe Phe
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 Gly Ala Val Leu Arg Arg Ala Ala Cys Leu Arg Arg Cys Leu Gly Pro
 115 120 125
 Pro Ser Ala His Leu Leu Ser Glu Glu Leu Asp Leu Glu Phe Asn Lys
 130 135 140
 Arg Ser Pro Tyr Asn Tyr Leu Gln Val Ala Tyr Phe Lys Ile Asn Lys
 145 150 155 160
 Leu Glu Lys Ala Val Ala Ala Ala His Thr Phe Phe Val Gly Asn Pro
 165 170 175
 Glu His Met Glu Met Arg Gln Asn Leu Asp Tyr Tyr Gln Thr Met Ser
 180 185 190
 Gly Val Lys Glu Ala Asp Phe Arg Asp Leu Glu Ala Lys Pro His Met
 195 200 205
 His Glu Phe Arg Leu Gly Val Arg Leu Tyr Ser Glu Glu Lys Pro Gln
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 Asp Glu Glu Cys Arg Ala Leu Cys Glu Gly Pro Tyr Asp Tyr Asp Gly

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Tyr Asn Tyr	Leu Asp Tyr Ser Ala Asp	Leu Phe Gln Ala Ile Thr Asp			
	260	265		270	
His Tyr Val	Gln Val Leu Asn Cys Lys Gln Asn Cys Val Thr Glu Leu				
	275	280		285	
Ala Ser His	Pro Ser Arg Glu Lys Pro Phe Glu Asp Phe Leu Pro Ser				
	290	295		300	
His Tyr Asn Tyr	Leu Gln Phe Ala Tyr Tyr Asn Ile Gly Asn Tyr Thr				
305	310	315		320	
Gln Ala Ile	Glu Cys Ala Lys Thr Tyr Leu Leu Phe Phe Pro Asn Asp				
	325	330		335	
Glu Val Met	His Gln Asn Leu Ala Tyr Tyr Thr Ala Met Leu Gly Glu				
	340	345		350	
Glu Glu Ala	Ser Ser Ile Ser Pro Arg Glu Asn Ala Glu Glu Tyr Arg				
	355	360		365	
Arg Pro Asn	Leu Leu Glu Lys Glu Leu Leu Phe Phe Ala Tyr Asp Ile				
	370	375		380	
Phe Gly Ile	Pro Phe Val Asp Pro Asp Ser Trp Thr Pro Glu Glu Val				
385	390	395		400	
Ile Pro Lys	Arg Leu Gln Glu Lys Gln Lys Ser Glu Arg Glu Thr Ala				
	405	410		415	
Val Arg Ile	Ser Gln Glu Ile Gly Asn Leu Met Lys Glu Ile Glu Thr				
	420	425		430	
Leu Val Glu	Lys Thr Lys Glu Ser Leu Asp Val Ser Arg Leu Thr				
	435	440		445	
Arg Glu Gly	Gly Pro Leu Leu Tyr Glu Gly Ile Ser Leu Thr Met Asn				
	450	455		460	
Ser Lys Val	Leu Asn Gly Ser Gln Arg Val Val Met Asp Gly Val Ile				
465	470	475		480	
Ser Asp Asp	Glu Cys Gln Glu Leu Gln Arg Leu Thr Asn Ala Ala Ala				
	485	490		495	
Thr Ser Gly	Asp Gly Tyr Arg Gly Gln Thr Ser Pro His Thr Pro Asn				
	500	505		510	
Glu Lys Phe	Tyr Gly Val Thr Val Leu Lys Ala Leu Lys Leu Gly Gln				
	515	520		525	
Glu Gly Lys	Val Pro Leu Gln Ser Ala Arg Thr Ala Leu Gln				
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